Figure 1

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241   gataaaaaccaaacgtacaacctcacggggctgcaaccttntacagaatatgtcatagct
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361   atgactgaggaagaangcaagctacttcctgcgattcccgtcctgtctgctctggtgtan
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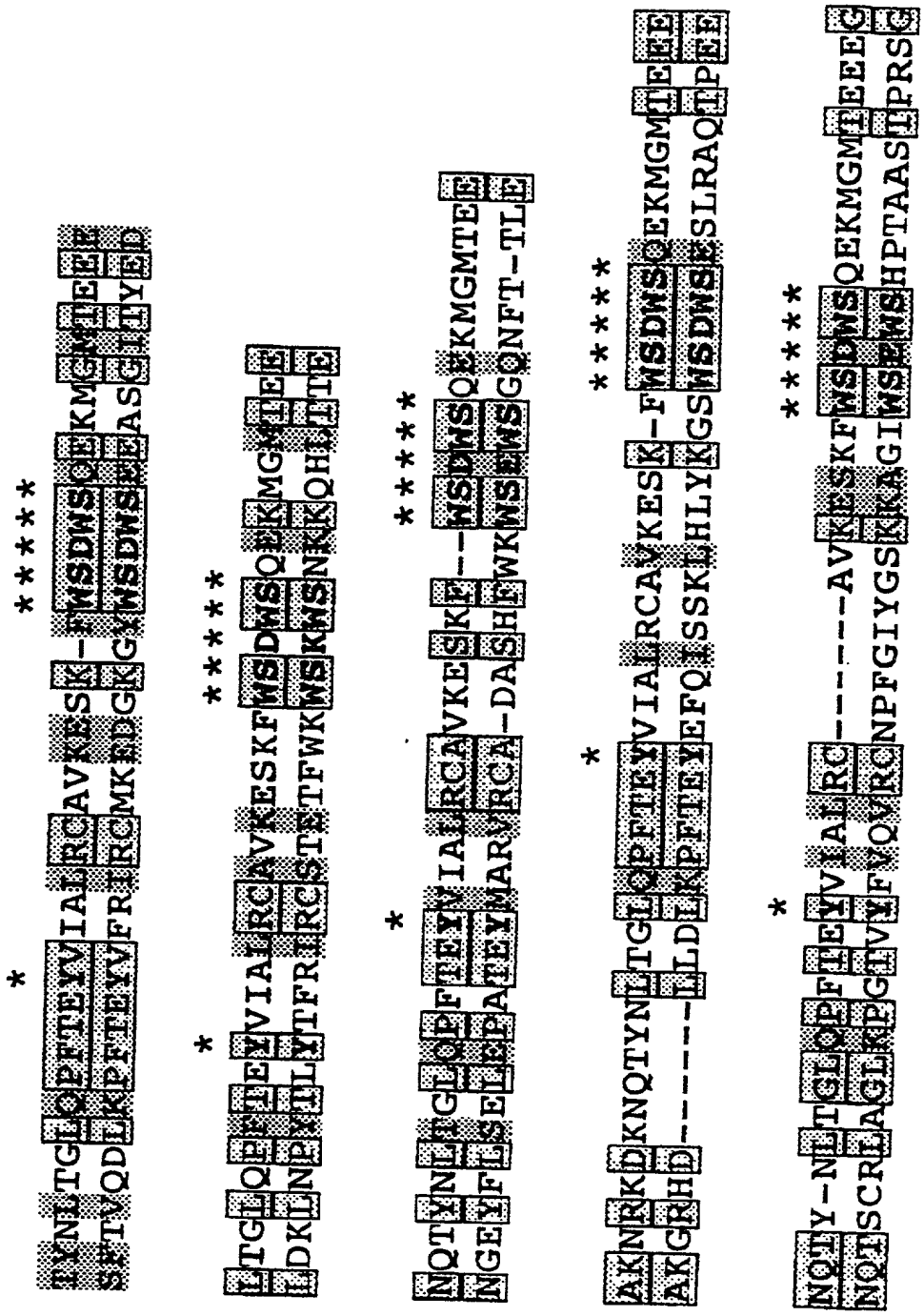
Figure 2



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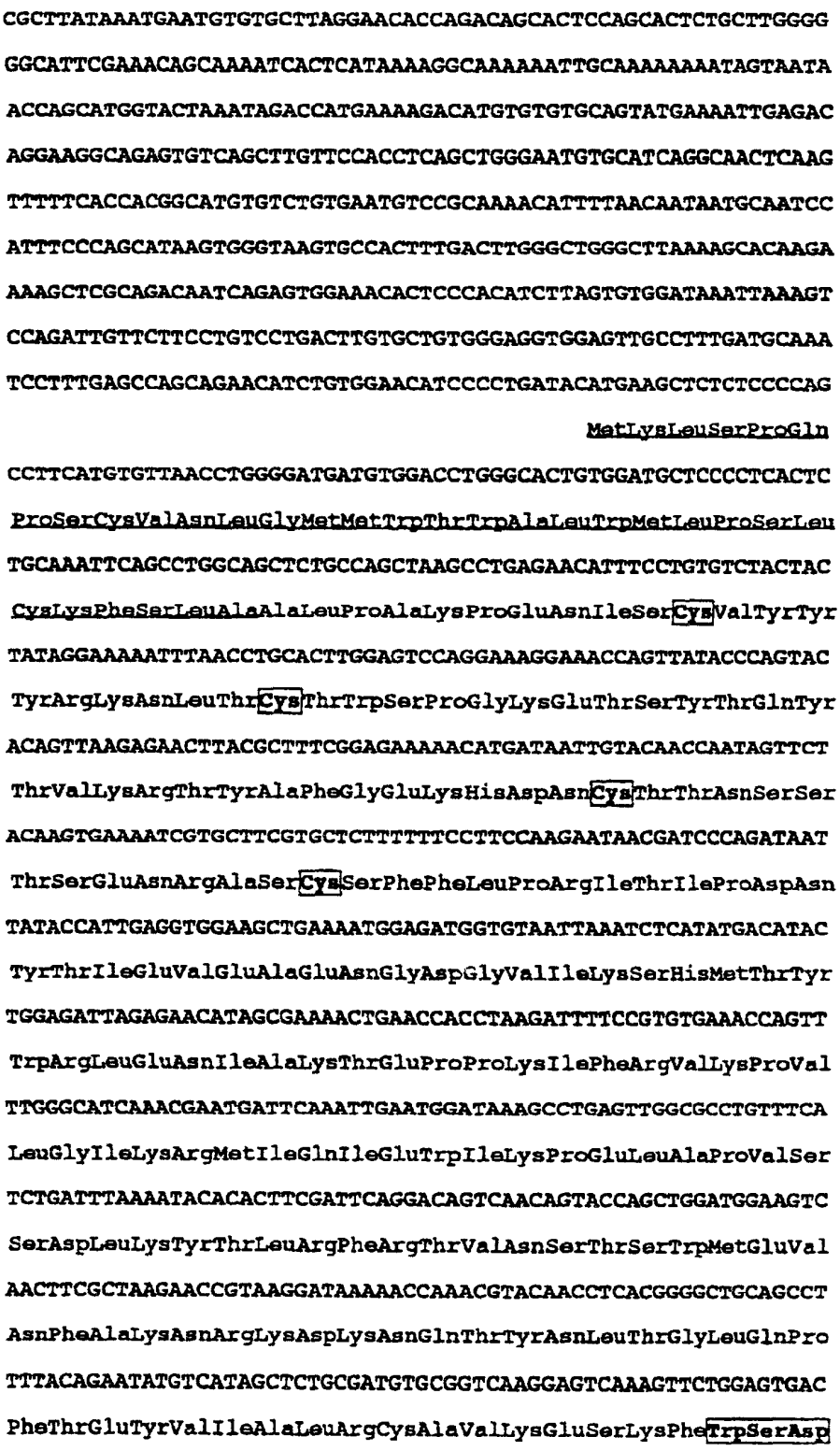
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241 TTTTTCACCACGGCATGTGTCTGTGAATGTCCGCAAAACATTTTAACAATAATGCAATCC
301 ATTTCCCAGCATAAGTGGGTAAGTGCCACTTTGACTTGGGCTGGGCTTAAAAGCACAAGA
361 AAAGCTCGCAGACAATCAGAGTGGAAACACTCCCACATCTTAGTGTGGATAAATTAAAGT
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481 TCCTTTGAGCCAGCAGAACATCTGTGGAACATCCCCTGATACATGAAGCTCTCTCCCCAG
MetLysLeuSerProGln
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ProSerCysValAsnLeuGlyMetMetTrpThrTrpAlaLeuTrpMetLeuProSerLeu
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Figure 4

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      ArgValLeuLysProAlaGluAlaAspGlyArgArgProValArgLeuLeuTrpLysLys
1321  GCAAGAGGAGCCCCAGTCCTAGAGAAAACACTTGGCTACAACATATGGTACTATCCAGAA
      AlaArgGlyAlaProValLeuGluLysThrLeuGlyTyrAsnIleTrpTyrTyrProGlu
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1861  GTGAAGACGGTCACGATCACATGGAAAGAGATTCCCAAGAGTGAGAGAAAGGGTATCATC
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      SerSerIleLeuGlnTyrGlyLeuGluSerLeuLysArgLysThrSerTyrIleValGln
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      ValMetAlaAsnThrSerAlaGlyGlyThrAsnGlyThrSerIleAsnPheLysThrLeu
2101  TCATTCAGTGTCTTTGAGATTATCCTCATAACTTCTCTGATTGGTGGAGGCCTTCTTATT
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Figure 5

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2281  AAGGATAAGCTAAACCTGAAGGAGTCTGATGACTCTGTGAACACAGAAGACAGGATCTTA
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Figure 6

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Figure 7

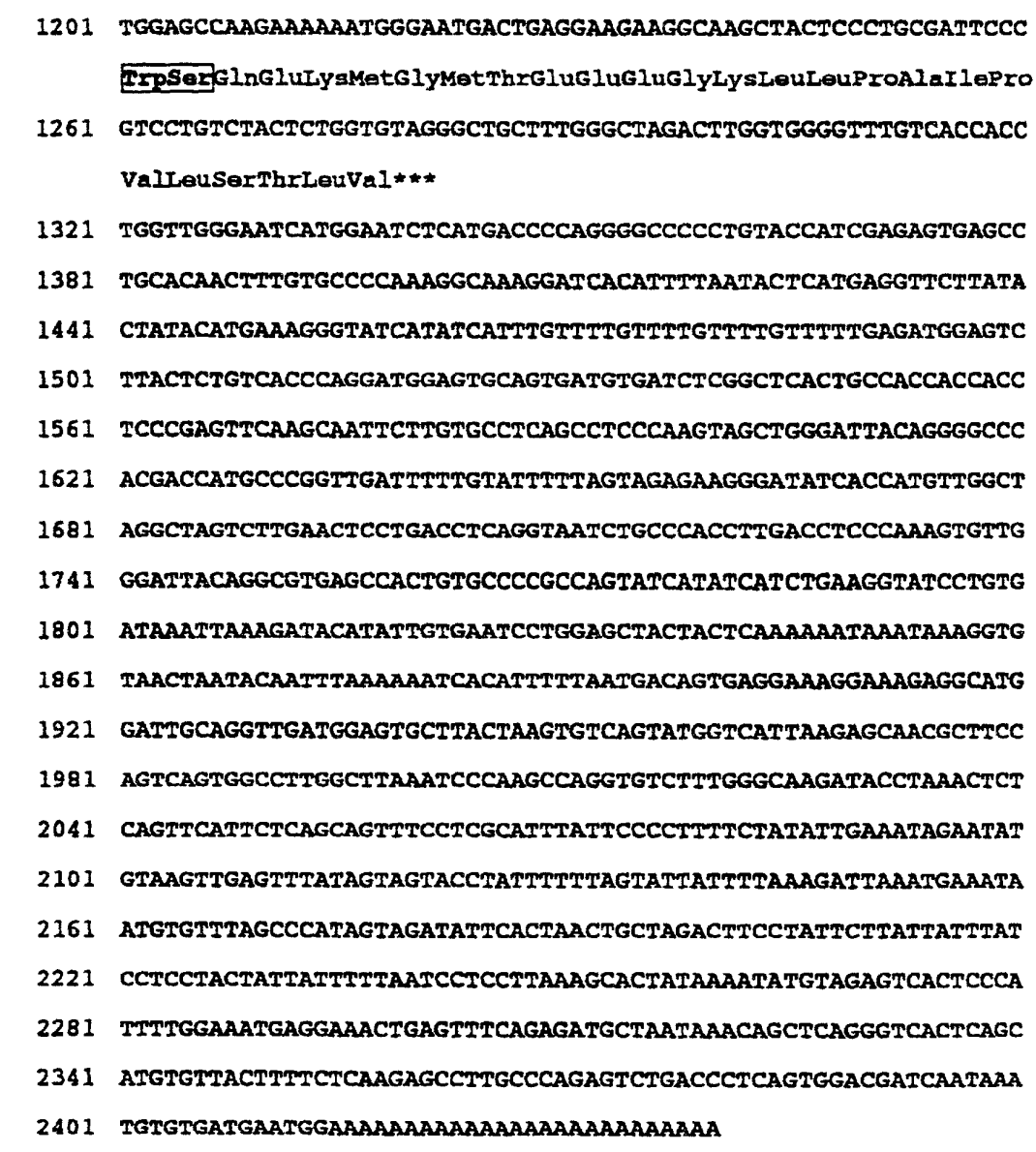
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Figure 8

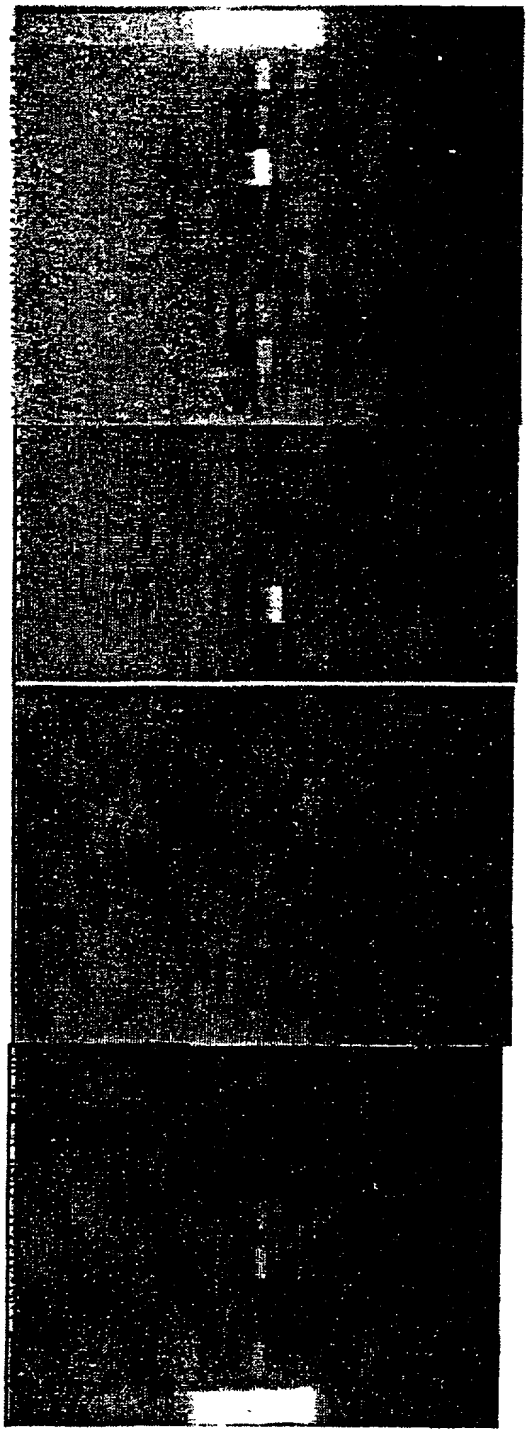
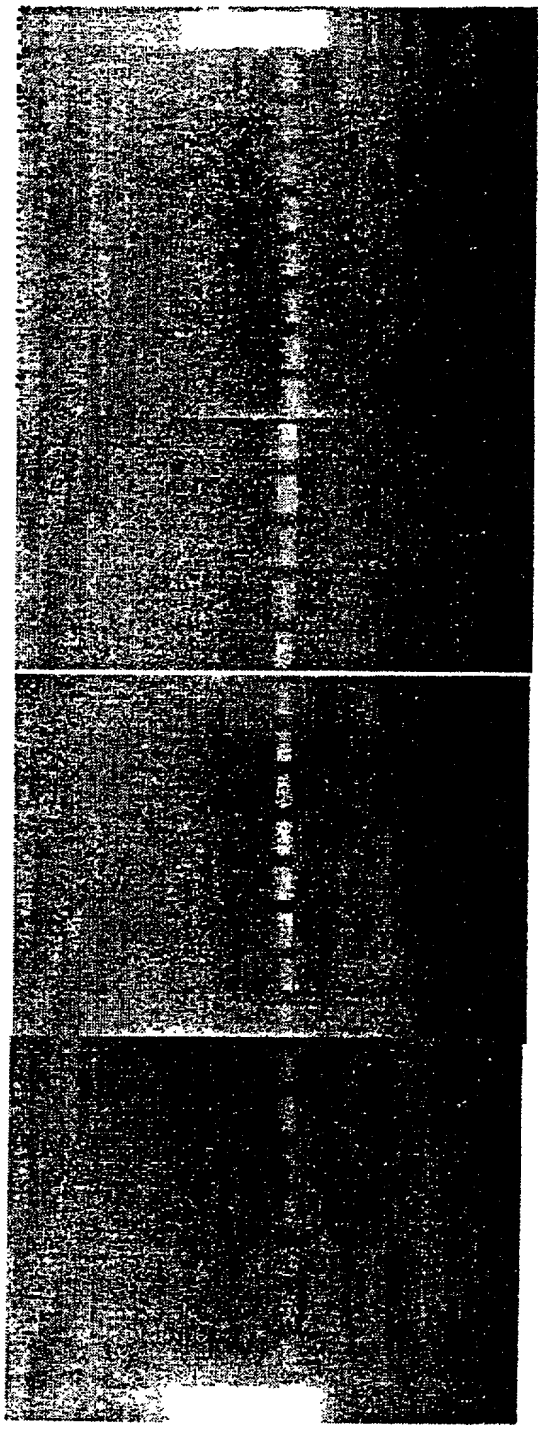


Figure 9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fetal spleen
fetal skeletal muscle
fetal lung
fetal liver
fetal spleen
fetal heart
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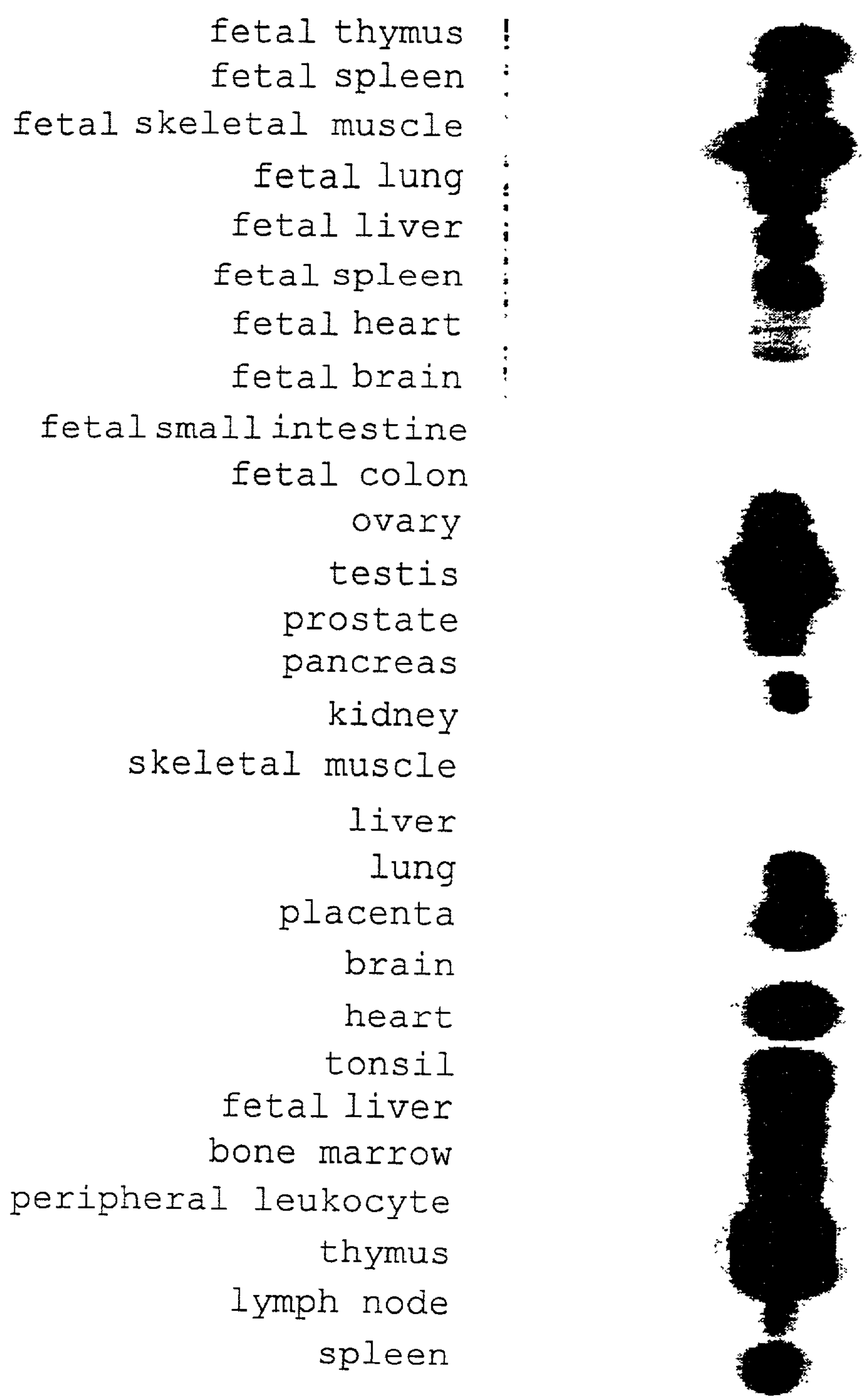
fetal small intestine
fetal colon
ovary
testis
prostate
pancreas
kidney
skeletal muscle
liver
lung
placenta
brain
heart
tonsil
fetal liver
bone marrow
peripheral leukocyte
thymus
lymph node
spleen

Figure 10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fetal spleen
fetal skeletal muscle
fetal lung
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fetal spleen
fetal heart
fetal brain
fetal small intestine
fetal colon
ovary
testis
prostate
pancreas
kidney
skeletal muscle
liver
lung
placenta
brain
heart
tonsil
fetal liver
bone marrow
peripheral leukocyte
thymus
lymph node
spleen

Figure 11

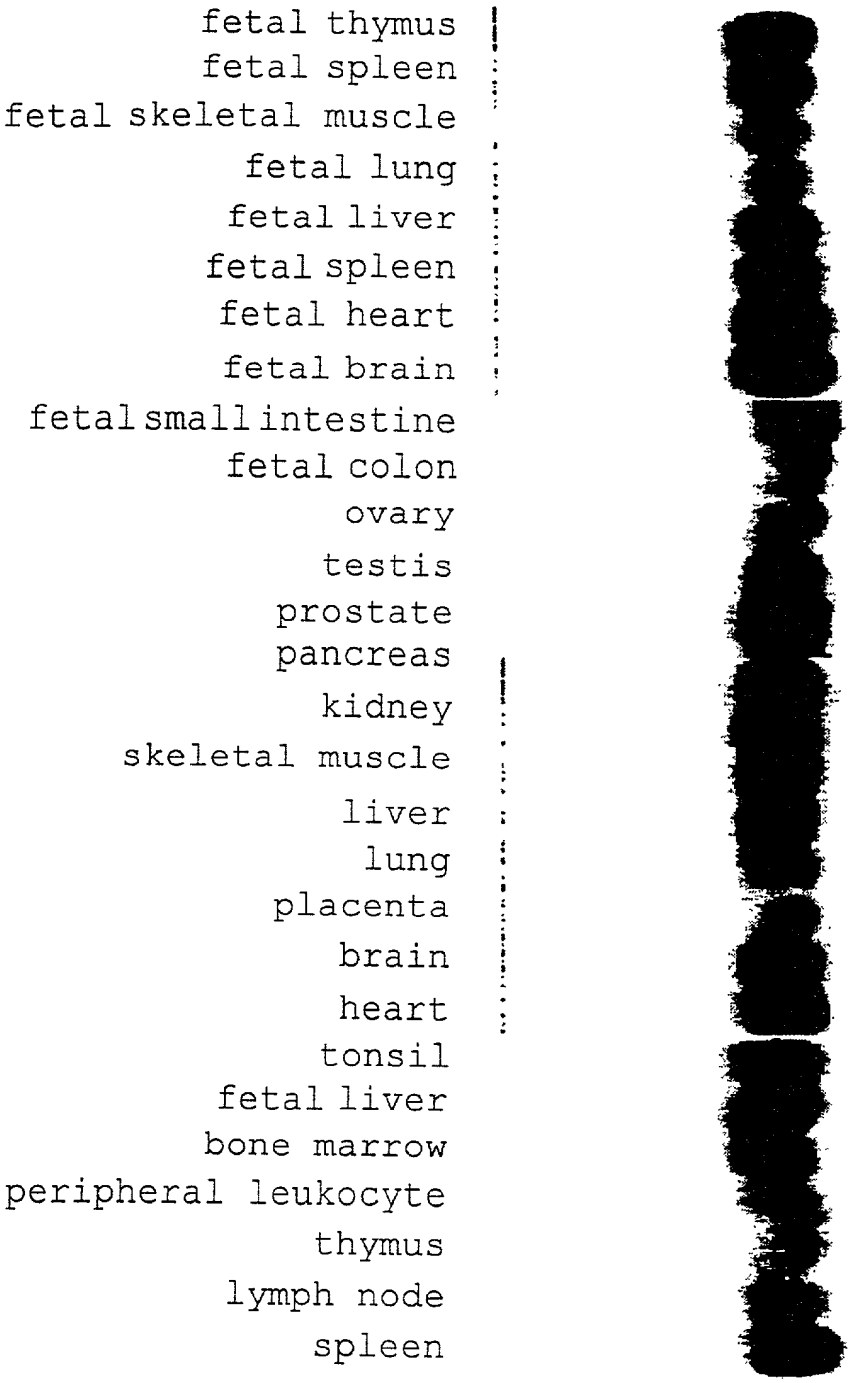

Figure 12

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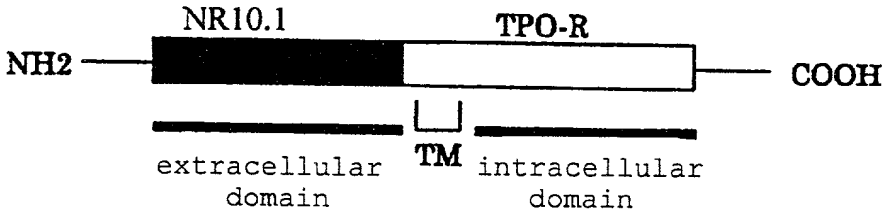

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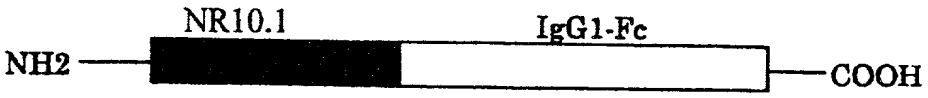

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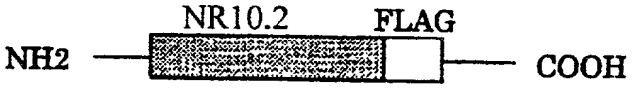

Figure 13

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Figure 14

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1741 TATAGCCACATGGCATGGAGATGATTTCAAGGATAAGCTAAACCTGAAGGAGTCTGATGA

IleAlaThrTrpHisGlyAspAspPheLysAspLysLeuAsnLeuLysGluSerAspAsp

1801 CTCTGTGAACACAGAAGACAGGATCTTAAAACCATGTTCCACCCCCAGTGACAAGTTGGT

SerValAsnThrGluAspArgIleLeuLysProCysSerThrProSerAspLysLeuVal

1861 GATTGACAAGTTGGTGGTGAACTTTGGGAATGTTCTGCAAGAAATTTTCACAGATGAAGC

IleAspLysLeuValValAsnPheGlyAsnValLeuGlnGluIlePheThrAspGluAla

1921 CAGAACGGGTCAGGAAAACAATTTAGGAGGGGAAAAGAATGGGACTAGAATTCTGTCTTC

ArgThrGlyGlnGluAsnAsnLeuGlyGlyGluLysAsnGlyThrArgIleLeuSerSer

1981 CTGCCCAACTTCAATATAAGTGTGGACTAAAATGCGAGAAAGGTGTCCTGTGGTCTATGC

CysProThrSerIle***

2041 AAATTAGAAAGGACATGCAGAGTTTTCCAACTAGGAAGACTGAATCTGTGGCCCCAAGAG

2101 AACCATCTCCGAAGACTGG